## Amendments to the Claims:

- 1-32. (cancelled)
- 33. (currently amended) An isolated nucleic acid comprising:
- (a) a nucleic acid sequence encoding the polypeptide shown in Figure 198 (SEQ ID NO:338);
- (b) a nucleic acid sequence encoding the polypeptide shown in Figure 198 (SEQ ID NO:338), lacking its associated signal peptide;
- (c) a nucleic acid sequence encoding the extracellular domain of the polypeptide shown in Figure 198 (SEQ ID NO:338);
- (d) a nucleic acid sequence encoding the extracellular domain of the polypeptide shown in Figure 198 (SEQ ID NO:338), lacking its associated signal peptide;
- [[(e)]](a) the nucleic acid sequence of SEQ ID NO:337 shown in Figure 197 (SEQ ID NO:337);
- [[(f)]](b) the full-length coding sequence of from within the nucleic acid sequence of SEQ ID NO:337shown in Figure 197 (SEQ ID NO:337); or
- [[(g)]] (c) the full-length coding sequence of the cDNA deposited under ATCC accession number 203322.
- 34. (currently amended) The isolated nucleic acid of Claim 33 comprising a nucleic acid sequence encoding the polypeptide of SEQ ID NO:338 shown in Figure 198 (SEQ ID NO:338).
- 35. (currently amended) The isolated nucleic acid of Claim 33 comprising a nucleic acid sequence encoding the polypeptide of SEQ ID NO:338 shown in Figure 198 (SEQ ID NO:338), lacking its associated signal peptide.
  - 36. (canceled)
  - 37. (canceled)

- 38. (currently amended) The isolated nucleic acid of Claim 33 comprising the nucleic acid sequence of SEQ ID NO:337 shown in Figure 197 (SEQ ID NO:337).
- 39. (currently amended) The isolated nucleic acid of Claim 33 comprising the full-length coding sequence of the nucleic acid sequence of SEQ ID NO:337 shown in Figure 197 (SEQ ID NO:337).
- 40. (previously presented) The isolated nucleic acid of Claim 33 comprising the full-length coding sequence of the cDNA deposited under ATCC accession number 203322.
  - 41. (canceled)
  - 42. (canceled)
  - 43. (canceled)
  - 44. (currently amended) A vector comprising the nucleic acid of Claim 28 33.
- 45. (previously presented) The vector of Claim 44, wherein said nucleic acid is operably linked to control sequences recognized by a host cell transformed with the vector.
  - 46. (previously presented) A host cell comprising the vector of Claim 44.
- 47. (previously presented) The host cell of Claim 46, wherein said cell is a CHO cell, an *E. coli* or a yeast cell.
- 48. (new) An isolated nucleic acid molecule at least 20 nucleotides in length that hybridizes under stringent conditions to:
  - (a) the nucleic acid sequence of SEQ ID NO:337 or a complement thereof;
- (b) the full-length coding sequence of the cDNA deposited under ATCC accession number 203322 or a complement thereof;

wherein, said stringent conditions use 50% formamide, 5 x SSC, 50 mM sodium phosphate (pH 6.8), 0.1% sodium pyrophosphate, 5x Denhardt's solution, sonicated salmon

sperm DNA (50  $\mu$ g/ml), 0.1% SDS, and 10% dextran sulfate at 42 °C, with washes at 42 °C in 0.2 x SSC and 50% formamide at 55 °C, followed by a wash comprising of 0.1 x SSC containing EDTA at 55 °C, wherein said isolated nucleic acid molecule is suitable for use as a PCR primer or probe.

- 49. (new) The isolated nucleic acid molecule of Claim 48 that is at least 50 nucleotides.
- 50. (new) The isolated nucleic acid molecule of Claim 48 that is at least 60 nucleotides.
- 51. (new) The isolated nucleic acid molecule of Claim 48 that is at least 70 nucleotides.
- 52. (new) The isolated nucleic acid molecule of Claim 48 that is at least 80 nucleotides.
- 53. (new) The isolated nucleic acid molecule of Claim 48 that is at least 90 nucleotides.
- 54. (new) The isolated nucleic acid molecule of Claim 48 that is at least 100 nucleotides.